Europäisches Patentamt European Patent Office Office européen des brevets



EP 0 990 047 B1 (11)

(12)

EUROPEAN PATENT SPECIFICATION

(45) Date of publication and mention of the grant of the patent:

14.05.2003 Bulletin 2003/20

(21) Application number: 98936928.5

(22) Date of filing: 22.07.1998

(51) Int CI.7: C12Q 1/68

(86) International application number: PCT/US98/15008

(87) International publication number: WO 99/005319 (04.02.1999 Gazette 1999/05)

(54) METHODS AND COMPOSITIONS FOR ANALYZING NUCLEIC ACIDS BY MASS SPECTROMETRY

VERFAHREN UND VERBINDUNGEN ZUR ANALYSIERUNG VON NUKLEINSAEUREN DURCH **MASSENSPEKTROMETRIE**

PROCEDES ET COMPOSITIONS POUR L'ANALYSE DE MOLECULES D'ACIDES NUCLEIQUES AU MOYEN DE SPECTROMETRIE DE MASSE

(84) Designated Contracting States: AT BE CH CY DE DK ES FI FR GB GR IE IT LI LU MC NL PT SE

(30) Priority: 22.07.1997 US 898180 22.07.1997 US 898564 22.07.1997 US 898501

(43) Date of publication of application: 05.04.2000 Bulletin 2000/14

(73) Proprietor: Qiagen Genomics, Inc. Bothell, Washington 98021 (US)

(72) Inventors:

 VAN NESS, Jeffrey Seattle, WA 98125 (US)

· TABONE, John, C. Bothell, WA 98011 (US)

 HOWBERT, Jeffry Bellevue, WA 98005 (US)

 MULLIGAN, John, T. Seattle, WA 98105 (US)

(74) Representative: Gowshall, Jonathan Vallance **FORRESTER & BOEHMERT** Pettenkoferstrasse 20-22 80336 München (DE)

(56) References cited:

WO-A-94/16101 WO-A-95/04160 WO-A-95/14108 WO-A-95/25737 WO-A-95/28640 WO-A-97/27325 WO-A-97/27327 WO-A-97/27331 US-A- 5 118 605

- JACOBSON K B ET AL: "APPLICATIONS OF MASS SPECTROMETRY TO DNA SEQUENCING" **GENETIC ANALYSIS TECHNIQUES AND** APPLICATIONS, vol. 8, no. 8, 1 December 1991, pages 223-229, XP000271820
- P.C.TORENET AL.: "Determination of Impurities in Nucleoside 3'-Phosphoramidites by Fast **Atom Bombardment Mass Spectrometry**" ANALYTICAL BIOCHEMISTRY, vol. 152, no. 2. 1986, pages 291-294, XP002094264
- **B.B.BROWN ET AL.: "A Single-Bead Decode** Strategy Using Electrospray Ionization Mass Spectometry and a New Photolabile Linker: 3-Amino-3-(2-nitrophenyl)Propionic Acid." MOLECULAR DIVERSITY, vol. 1, 1995, pages 4-12, XP002094265

Note: Within nine months from the publication of the mention of the grant of the European patent, any person may give notice to the European Patent Office of opposition to the European patent granted. Notice of opposition shall be filed in a written reasoned statement. It shall not be deemed to have been filed until the opposition fee has been paid. (Art. 99(1) European Patent Convention).

Claims

1. A compound of the formula:

T^{ms}-L-X

wherein,

T^{ms} is an organic group detectable by mass spectrometry, comprising carbon, at least one of hydrogen and fluoride, and optional atoms selected from oxygen, nitrogen, sulfur, phosphorus and iodine;

L is an organic group which allows a unique T^{ms}-containing moiety to be cleaved from the remainder of the compound, wherein the T^{ms}-containing moiety comprises a functional group which supports a single ionized charge state when the compound is subjected to mass spectrometry and is tertiary amine, quaternary amine or organic acid;

X is a functional group selected from phosphoramidite and H-phosphonate.

2. The compound of claim 1 wherein X is a phosphoramidite group such that Tms-L-X has the structure

20

5

10

15

25

wherein R is an alkyl group or a substituted alkyl group having one or more substituents selected from halogen and cyano, and the two R groups of NR₂ may be bonded together to form a cycloalkyl group.

3. The compound of claim 2 wherein X is a phosphoramidite group such that Tms-L-X has the structure

35

40

45

30

and OR is OCH2CH2CN while NR2 is N(iso-propyl)2.

4. The compound of claim 1 wherein X has an H-phosphonate group such that T^{ms}-L-X has the structure

50

wherein R is a C₁-C₆alkyl group.

55

5. A compound according to claims 1-4 wherein T^{ms} has a mass of from 15 to 10,000 daltons and a molecular formula of $C_{1-500}N_{0-100}O_{0-100}S_{0-10}P_{0-10}H_{\alpha}F_{\beta}I_{\delta}$ wherein the sum of α , β and δ is sufficient to satisfy the otherwise unsatisfied valencies of the C, N and O atoms.

55

6. A compound according to claims 1-4 wherein T^{ms} and L are bonded together through a functional group selected from amide, ester, ether, amine, sulfide, thioester, disulfide, thioether, urea, thiourea, carbamate, thiocarbamate, Schiff base, reduced Schiff base, imine, oxime, hydrazone, phosphate, phosphonate, phosphoramide, phosphona-

mide, sulfonate, sulfonamide or carbon-carbon bond.

5

10

15

20

35

40

50

- 7. A compound according to claims 1-4 wherein L is selected from L^{hv}, L^{acid}, L^{base}, L^(O), L^(R), L^{enz}, L^{elc}, L^Δ and L^{ss}, where actinic radiation, acid, base, oxidation, reduction, enzyme, electrochemical, thermal and thiol exchange, respectively, cause the T^{ms}-containing moiety to be cleaved from the remainder of the molecule.
- 8. A compound according to claims 1-4 wherein Lhv has the formula L1-L2-L3, wherein L2 is a molecular fragment that absorbs actinic radiation to promote the cleavage of Tms from X, and L1 and L3 are independently a direct bond or an organic moiety, where L1 separates L2 from Tms and L3 separates L2 from X, and neither L1 nor L3 undergo bond cleavage when L2 absorbs the actinic radiation.
- 9. A compound according to claim 8 wherein -L2-L3 has the formula:

- with one carbon atom at positions a, b, c, d or e being substituted with -L³-X and optionally one or more of positions b, c, d or e being substituted with alkyl, alkoxy, fluoride, chloride, hydroxyl, carboxylate or amide; and R¹ is hydrogen or hydrocarbyl.
- 10. A compound according to claim 8 wherein L³ is selected from a direct bond, a hydrocarbylene, -O-hydrocarbylene, and hydrocarbylene-(O-hydrocarbylene)_n-H, and n is an integer ranging from 1 to 10.
 - 11. A compound according to claims 1-4 wherein -L-X has the formula:

$$\begin{array}{c}
c & d \\
b & NO_2 \\
-N & C-R^2 \\
0
\end{array}$$

- wherein one or more of positions b, c, d or e is substituted with hydrogen, alkyl, alkoxy, fluoride, chloride, hydroxyl, carboxylate or amide; R¹ is hydrogen or hydrocarbyl, and R² terminates in an "X" group.
 - 12. A compound according to claims 1-4 wherein Tms has the formula:

$$T^2$$
-(J- T^3 -)_n-

- T² is an organic moiety formed from carbon and one or more of hydrogen, fluoride, iodide, oxygen, nitrogen, sulfur and phosphorous, having a mass of 15 to 500 daltons;
- T³ is an organic molety formed from carbon and one or more of hydrogen, fluoride, lodide, oxygen, nitrogen, sulfur and phosphorous, having a mass of 50 to 1000 daltons;
- J is a direct bond or a functional group selected from amide, ester, amine, sulfide, ether, thioester, disulfide, thioether, urea, thiourea, carbamate, thiocarbamate, Schiff base, reduced Schiff base, imine, oxime, hydra-

zone, phosphate, phosphonate, phosphoramide, phosphonamide, sulfonate, sulfonamide or carbon-carbon

n is an integer ranging from 1 to 50, and when n is greater than 1, each T3 and J is independently selected.

- 13. A compound according to claim 12 wherein T² is selected from hydrocarbyl, hydrocarbyl-O-hydrocarbylene, hydrocarbyl-S-hydrocarbylene, hydrocarbyl-NH-hydrocarbylene, hydrocarbyl-amide-hydrocarbylene, N-(hydrocarbyl)hydrocarbylene, N,N-di(hydrocarbyl)hydrocarbylene, hydrocarbylacyl-hydrocarbylene, heterocyclylhydrocarbyl wherein the heteroatom(s) are selected from oxygen, nitrogen, sulfur and phosphorous, substituted heterocyclylhydrocarbyl wherein the heteroatom(s) are selected from oxygen, nitrogen, sulfur and phosphorous and the sub-10 stituents are selected from hydrocarbyl, hydrocarbyl-O-hydrocarbylene, hydrocarbyl-NH-hydrocarbylene, hydrocarbyl-S-hydrocarbylene, N-(hydrocarbyl)hydrocarbylene, N,N-di(hydrocarbyl)hydrocarbylene and hydrocarbylacyl-hydrocarbylene, as well as derivatives of any of the foregoing wherein one or more hydrogens is replaced with an equal number of fluorides.
- 15 14. A compound according to claim 12 wherein T^3 has the formula - $G(R^2)$ -, G is C_{1-6} alkylene having a single R^2 substituent, and R2 is selected from alkyl, alkenyl, alkynyl, cycloalkyl, aryl-fused cycloalkyl, cycloalkenyl, aryl, aralkyl, aryl-substituted alkenyl or alkynyl, cycloalkyl-substituted alkyl, cycloalkenyl-substituted cycloalkyl, biaryl, alkoxy, alkenoxy, alkynoxy, aralkoxy, aryl-substituted alkenoxy or alkynoxy, alkylamino, alkenylamino or alkynylamino, aryl-substituted alkylamino, aryl-substituted alkenylamino or alkynylamino, aryloxy, arylamino, N-alkylurea-20 substituted alkyl, N-arylurea-substituted alkyl, alkylcarbonylamino-substituted alkyl, aminocarbonyl-substituted alkyl, heterocyclyl, heterocyclyl-substituted alkyl, heterocyclyl-substituted amino, carboxyalkyl substituted aralkyl, oxocarbocyclyl-fused aryl and heterocyclylalkyl; cycloalkenyl, aryl-substituted alkyl and, aralkyl, hydroxy-substituted alkyl, alkoxy-substituted alkyl, aralkoxy-substituted alkyl, alkoxy-substituted alkyl, aralkoxy-substituted alkyl, amino-substituted alkyl, (aryl-substituted alkyloxycarbonylamino)-substituted alkyl, thiol-substituted alkyl, alkylsul-25 fonyl-substituted alkyl, (hydroxy-substituted alkylthio)-substituted alkyl, thioalkoxy-substituted alkyl, hydrocarbylacylamino-substituted alkyl, heterocyclylacylamino-substituted alkyl, hydrocarbyl-substituted-heterocyclylacylamino-substituted alkyl, alkylsulfonylamino-substituted alkyl, arylsulfonylamino-substituted alkyl, morpholino-alkyl, thiomorpholino-alkyl, morpholino carbonyl-substituted alkyl, thiomorpholinocarbonyl-substituted alkyl, [N-(alkyl, alkenyl or alkynyl)- or N,N-[dialkyl, dialkenyl, dialkynyl or (alkyl, alkenyl)-amino]carbonyl-substituted alkyl, hetero-30 cyclylaminocarbonyl, heterocylylalkyleneaminocarbonyl, heterocyclylaminocarbonyl-substituted alkyl, heterocylylalkyleneaminocarbonyl-substituted alkyl, N,N-[dialkyl]alkyleneaminocarbonyl, N,N-[dialkyl]alkyleneaminocarbonyl-substituted alkyl, alkyl-substituted heterocyclylcarbonyl, alkyl-substituted heterocyclylcarbonyl-alkyl, carboxylsubstituted alkyl, dialkylamino-substituted acylaminoalkyl and amino acid side chains selected from arginine, asparagine, glutamine, S-methyl cysteine, methionine and corresponding sulfoxide and sulfone derivatives thereof, 35 glycine, leucine, isoleucine, allo-isoleucine, tert-leucine, norteucine, phenylalanine, tyrosine, tryptophan, proline, alanine, ornithine, histidine, glutamine, valine, threonine, serine, aspartic acid, beta-cyanoalanine, and allothreonine; alynyl and heterocyclylcarbonyl, aminocarbonyl, amido, mono- or dialkylaminocarbonyl, mono- or diarylaminocarbonyl, alkylarylaminocarbonyl, diarylaminocarbonyl, mono- or diacylaminocarbonyl, aromatic or aliphatic acyl, alkyl optionally substituted by substituents selected from amino, carboxy, hydroxy, mercapto, mono- or dialkylamino, mono- or diarylamino, alkylarylamino, diarylamino, mono- or diacylamino, alkoxy, alkenoxy, aryloxy, 40 thioalkoxy, thioalkenoxy, thioalkynoxy, thioaryloxy and heterocyclyl.
 - 15. A compound according to claim 12 having the formula:

$$T^{4}$$
Amide
$$(CH_{2})_{c}$$

$$T^{2}$$

$$N$$

$$R^{1}$$

$$X$$

wherein

5

45

50

G is $(CH_2)_{1-6}$ wherein a hydrogen on one and only one of the CH_2 groups is replaced with- $(CH_2)_c$ -Amide- T^4 ; T^2 and T^4 are organic moieties of the formula $C_{1-25}N_{0-9}O_{0-9}H_{\alpha}F_{\beta}$ wherein the sum of α and β is sufficient to satisfy the otherwise unsatisfied valencies of the C, N, and O atoms;

 R^1 is hydrogen or C_{1-10} alkyl; c is an integer ranging from 0 to 4;

5

10

15

20

25

30

35

40

45

50

55

X is defined according to claim 1; and

n is an integer ranging from 1 to 50 such that when n is greater than 1, G, c, Amide, R^1 and T^4 are independently selected.

16. A compound according to claim 15 having the formula:

Amide

$$CH_2$$
 CH_2
 CH_2

wherein T⁵ is an organic moiety of the formula $C_{1-25}N_{0-9}O_{0-9}H_{\alpha}F_{\beta}$ wherein the sum of α and β is sufficient to satisfy the otherwise unsatisfied valencies of the C, N, and O atoms; and T⁵ includes a tertiary or quaternary amine or an organic acid; and m is an integer ranging from 0-49.

17. A compound according to claim 15 having the formula:

$$T^{2} \xrightarrow{\text{Amide}} L^{X}$$

$$T^{2} \xrightarrow{\text{R}^{1}} O$$

$$R^{1} \xrightarrow{\text{Amide}} L^{X}$$

$$T^{3} \xrightarrow{\text{Amide}} L^{X}$$

wherein T⁵ is an organic moiety of the formula $C_{1-25}N_{0-9}O_{0-9}H_{\alpha}F_{\beta}$ wherein the sum of α and β is sufficient to satisfy the otherwise unsatisfied valencies of the C, N, and O atoms; and T⁵ includes a tertiary or quaternary

amine or an organic acid; and m is an integer ranging from 0-49.

18. A compound according to any one of claims 16 and 17 wherein -Amide-T⁵ is selected from:

$$-NHC \longrightarrow N ; -NHC \longrightarrow O - (C_2 - C_{10}) - N(C_1 - C_{10})_2$$

$$(C_1 - C_{10})$$

$$-NHC - (C_1 - C_{10}) - N \longrightarrow : -NHC \longrightarrow N ;$$

$$-NHC \longrightarrow N-(C_1-C_{10});$$
 and $-NHC-(C_1-C_{10})-N \longrightarrow 0$

19. A compound according to any of claims 16 and 17 wherein -Amide-T⁵ is selected from:

$$-\frac{\text{CNH}-(C_1-C_{10})}{0}; \quad -\frac{\text{CNH}-(C_2-C_{10})}{0};$$

$$- \begin{array}{c} (C_1 - C_{10}) \\ - (C_1 - C_{10}) - (C_1 - C_{10}) \\ 0 \end{array}; \quad - \begin{array}{c} (C_1 - C_{10}) \\ - (C_1 - C_{10}) - (C_1 - C_{10}) \\ 0 \end{array}$$

$$-CNH - (C_2 - C_{10}) - N(C_1 - C_{10})_2; \quad -CNH - (C_2 - C_{10}) - N$$
;

$$-CN N(C_1-C_{10}); \text{ and } CN NH N$$

20. A compound according to any one of claims 14-16 wherein T² has the structure which results when one of the following organic acids is condensed with an amine group to form T²-C(=O)-N(R¹)-: Formic acid, Acetic acid, Propiolic acid, Propiolic acid, Fluoroacetic acid, 2-Butynoic acid, Cyclopropanecarboxylic acid, Butyric acid, Methoxyacetic acid, Difluoroacetic acid, 4-Pentynoic acid, Cyclobutanecarboxylic acid, 3,3-Dimethylacrylic acid, Valeric

acid, N,N-Dimethylglycine, N-Formyl-Gly-OH, Ethoxyacetic acid, (Methylthio)acetic acid, Pyrrole-2-carboxylic acid, 3-Furoic acid, Isoxazole-5-carboxylic acid, trans-3-Hexenoic acid, Trifluoroacetic acid, Hexanoic acid, Ac-Gly-OH, 2-Hydroxy-2-methylbutyric acid, Benzoic acid, Nicotinic acid, 2-Pyrazinecarboxylic acid, 1-Methyl-2-pyrrolecarboxylic acid, 2-Cyclopentene-1-acetic acid, Cyclopentylacetic acid, (S)-(-)-2-Pyrrolidone-5-carboxylic acid, N-Methyl-L-proline, Heptanoic acid, Ac-b-Ala-OH, 2-Ethyl-2-hydroxybutyric acid, 2-(2-Methoxyethoxy)acetic acid, p-Toluic acid, 6-Methylnicotinic acid, 5-Methyl-2-pyrazinecarboxylic acid, 2,5-Dimethylpyrrole-3-carboxylic acid, 4-Fluorobenzoic acid, 3,5-Dimethylisoxazole-4-carboxylic acid, 3-Cyclopentylpropionic acid, Octanoic acid, N, N-Dimethylsuccinamic acid, Phenylpropiolic acid, Cinnamic acid, 4-Ethylbenzoic acid, p-Anisic acid, 1,2,5-Trimethylpyrrole-3-carboxylic acid, 3-Fluoro-4-methylbenzoic acid, Ac-DL-Propargylglycine, 3-(Trifluoromethyl)butyric acid, 1-Piperidinepropionic acid, N-Acetylproline, 3,5-Difluorobenzoic acid, Ac-L-Val-OH, Indole-2-carboxylic acid, 2-Benzofurancarboxylic acid, Benzotriazole-5-carboxylic acid, 4-n-Propylbenzoic acid, 3-Dimethylaminobenzoic acid, 4-Ethoxybenzoic acid, 4-(Methylthio)benzoic acid, N-(2-Furoyl)glycine, 2-(Methylthio)nicotinic acid, 3-Fluoro-4-methoxybenzoic acid, Tfa-Gly-OH, 2-Napthoic acid, Quinaldic acid, Ac-L-Ile-OH, 3-Methylindene-2-carboxylic acid, 2-Quinoxalinecarboxylic acid, 1-Methylindole-2-carboxylic acid, 2,3,6-Trifluorobenzoic acid, N-Formyl-L-Met-OH, 2-[2-(2-Methoxyethoxy)ethoxy]acetic acid, 4-n-Butylbenzoic acid, N-Benzoylglycine, 5-Fluoroindole-2-carboxylic acid, 4-n-Propoxybenzoic acid, 4-Acetyl-3,5-dimethyl-2-pyrrolecarboxylic acid, 3,5-Dimethoxybenzoic acid, 2,6-Dimethoxynicotinic acid, Cyclohexanepentanoic acid, 2-Naphthylacetic acid, 4-(1H-Pyrrol-1-yl)benzoic acid, Indole-3-propionic acid, m-Trifluoromethylbenzoic acid, 5-Methoxyindole-2-carboxylic acid, 4-Pentylbenzoic acid, Bz-b-Ala-OH, 4-Diethylaminobenzoic acid, 4-n-Butoxybenzoic acid, 3-Methyl-5-CF3-isoxazole-4-carboxylic acid, (3,4-Dimethoxyphenyl)acetic acid, 4-Biphenylcarboxylic acid, Pivaloyl-Pro-OH, Octanoyl-Gly-OH, (2-Naphthoxy)acetic acid, Indole-3-butyric acid, 4-(Trifluoromethyl)phenylacetic acid, 5-Methoxyindole-3-acetic acid, 4-(Trifluoromethoxy)benzoic acid, Ac-L-Phe-OH, 4-Pentyloxybenzoic acid, Z-Gly-OH, 4-Carboxy-N-(fur-2-ylmethyl)pyrrolidin-2-one, 3,4-Diethoxybenzoic acid, 2,4-Dimethyl-5-CO₂Et-pyrrole-3-carboxylic acid, N-(2-Fluorophenyl)succinamic acid, 3,4,5-Trimethoxybenzoic acid, N-Phenylanthranilic acid, 3-Phenoxybenzoic acid, Nonanoyl-Gly-OH, 2-Phenoxypyridine-3-carboxylic acid, 2,5-Dimethyl-1-phenylpyrrole-3-carboxylic acid, trans-4-(Trifluoromethyl) cinnamic acid, (5-Methyl-2-phenyloxazol-4-yl)acetic acid, 4-(2-Cyclohexenyloxy)benzoic acid, 5-Methoxy-2-methylindole-3-acetic acid, trans-4-Cotininecarboxylic acid, Bz-5-Aminovaleric acid, 4-Hexyloxybenzoic acid, N-(3-Methoxyphenyl)succinamic acid, Z-Sar-OH, 4-(3,4-Dimethoxyphenyl)butyric acid, Ac-o-Fluoro-DL-Phe-OH, N-(4-Fluorophenyl)glutaramic acid, 4'-Ethyl-4-biphenylcarboxylic acid, 1,2,3,4-Tetrahydroacridinecarboxylic acid, 3-Phenoxyphenylacetic acid, N-(2,4-Difluorophenyl)succinamic acid, N-Decanoyl-Gly-OH, (+)-6-Methoxy-a-methyl-2-naphthaleneacetic acid, 3-(Trifluoromethoxy)cinnamic acid, N-Formyl-DL-Trp-OH, (R)-(+)-a-Methoxy-a-(trifluoromethyl)phenylacetic acid, Bz-DL-Leu-OH, 4-(Trifluoromethoxy)phenoxyacetic acid, 4-Heptyloxybenzoic acid, 2,3,4-Trimethoxycinnamic acid, 2,6-Dimethoxybenzoyl-Gly-OH, 3-(3,4,5-Trimethoxyphenyl)propionic acid, 2,3,4,5,6-Pentafluorophenoxyacetic acid, N-(2,4-Difluorophenyl)glutaramic acid, N-Undecanoyl-Gly-OH, 2-(4-Fluorobenzoyl)benzoic acid, 5-Trifluoromethoxyindole-2-carboxylic acid, N-(2,4-Difluorophenyl)diglycolamic acid, Ac-L-Trp-OH, Tfa-L-Phenylglycine-OH, 3-lodobenzoic acid, 3-(4-n-Pentylbenzoyl)propionic acid, 2-Phenyl-4-quinolinecarboxylic acid, 4-Octyloxybenzoic acid, Bz-L-Met-OH, 3,4,5-Triethoxybenzoic acid, N-Lauroyl-Gly-OH, 3,5-Bis(trifluoromethyl)benzoic acid, Ac-5-Methyl-DL-Trp-OH, 2-lodophenylacetic acid, 3-lodo-4-methylbenzoic acid, 3-(4-n-Hexylbenzoyl)propionic acid, N-Hexanoyl-L-Phe-OH, 4-Nonyloxybenzoic acid, 4'-(Trifluoromethyl)-2-biphenylcarboxylic acid, Bz-L-Phe-OH, N-Tridecanoyl-Gly-OH, 3,5-Bis(trifluoromethyl)phenylacetic acid, 3-(4-n-Heptylbenzoyl)propionic acid, N-Hepytanoyl-L-Phe-OH, 4-Decyloxybenzoic acid, N- $(\alpha,\alpha,\alpha-trifluoro-m-tolyl)$ anthranilic acid, Niflumic acid, 4-(2-Hydroxyhexafluoroisopropyl)benzoic acid, N-Myristoyl-Gly-OH, 3-(4-n-Octylbenzoyl)propionic acid, N-Octanoyl-L-Phe-OH, 4-Undecyloxybenzoic acid, 3-(3,4,5-Trimethoxyphenyl)propionyl-Gly-OH, 8-lodonaphthoic acid, N-Pentadecanoyl-Gly-OH, 4-Dodecyloxybenzoic acid, N-Palmitoyl-Gly-OH, and N-Stearovi-Gly-OH.

- 21. A method for determining the presence of a single nucleotide polymorphism in a nucleic acid target comprising:
 - a) amplifying a sequence of a nucleic acid target containing a single nucleotide polymorphism;
 - b) generating a single strand form of the target;

5

10

· 15

20

25

30

35

40

45

50

- c) combining a tagged nucleic acid probe with the amplified target nucleic acid molecules under conditions and for a time sufficient to permit hybridization of said tagged nucleic acid probe to complementary amplified selected target nucleic acid molecules, wherein said tag is correlative with a particular single nucleotide polymorphism and is detectable by spectrometry or potentiometry;
- d) separating unhybridized tagged probe from hybridized tagged probe by a sizing methodology;
- e) cleaving said tag from said probe; and
- f) detecting said tag by spectrometry or potentiometry, and determining the presence of said single nucleotide polymorphism,

wherein the tagged nucleic acid probe has the structure

T^{ms}-L-X'

5

10

wherein.

T^{ms} is an organic group detectable by mass spectrometry, comprising carbon, at least one of hydrogen and fluoride, and optional atoms selected from oxygen, nitrogen, sulfur, phosphorus and iodine;

L is an organic group which allows a unique T^{ms}-containing moiety to be cleaved from the remainder of the compound, wherein the T^{ms}-containing moiety, comprises a functional group which supports a single ionized charge state when the compound is subjected to mass spectrometry and is tertiary amine, quaternary amine or organic acid; and

X' is nucleic acid.

15

20

25

- 22. A method for determining the presence of a single nucleotide polymorphism in a nucleic acid target comprising:
 - a) amplifying a sequence of a nucleic acid target containing a single nucleotide polymorphism:
 - b) combining a tagged nucleic acid primer with the amplified target nucleic acid molecules under conditions and for a time sufficient to permit annealing of said tagged nucleic acid primer to complementary amplified selected target nucleic acid molecules, wherein the oligonucleotide primer has a 3'-most base complementary to the wildtype sequence or the single nucleotide polymorphism, wherein said tag is correlative with a particular single nucleotide polymorphism and is detectable by spectrometry or potentiometry;
 - c) extending the primer wherein a complementary strand to the target is synthesized when the 3'-most baseof the primer is complementary to the target;
 - d) separating unextended tagged primer from extended tagged primer by a sizing methodology;
 - e) cleaving said tag from said primers or extended primers; and
 - f) detecting said tag by spectrometry or potentiometry, and determining therefrom the presence of said single nucleotide polymorphis,

30

wherein the tagged nucleic acid primer has the structure

T^{ms}-L-X'

35

40

wherein,

T^{ms} is an organic group detectable by mass spectrometry, comprising carbon, at least one of hydrogen and fluoride, and optional atoms selected from oxygen, nitrogen, sulfur, phosphorus and iodine;

L is an organic group which allows a unique Tms-containing moiety to be cleaved from the remainder of the compound, wherein the Tms-containing moiety comprises a functional group which supports a single ionized charge state when the compound is subjected to mass spectrometry and is tertiary amine, quaternary amine or organic acid; and

X' is nucleic acid.

45

50

- 23. A method for determining the quantity of a specific mRNA molecule in a nucleic acid population comprising:
 - a) converting an RNA population into a cDNA population;
 - b) adding a single strand nucleic acid (internal standard) containing a plurality of single nucleotide polymorphisms, that is otherwise identical to said cDNA target;
 - c) amplifying a specific sequence of said cDNA target;
 - d) coamplifying the internal standard, wherein said internal standard is the same length as the cDNA amplicon;
 - e) generating a single strand form of the target;
 - f) combining a set of tagged nucleic acid probes with the amplified target cDNA and amplified internal standard under conditions and for a time sufficient to permit hybridization of said tagged nucleic acid probe to complementary selected target cDNA and internal standard sequences, wherein said tag is correlative with a particular cDNA sequence and a second tag is correlative with the internal standard, and is detectable by spectrometry or potentiometry;

- g) separating unhybridized tagged probe from hybridized tagged probe by a sizing methodology;
- h) cleaving said tag from said probes;
- i) detecting said tags by spectrometry or potentiometry; and
- j) taking the ratio of tag correlated to cDNA to tag correlated with the internal standard, and determining therefrom the quantity of said cDNA, thereby determining the quantity of the specific mRNA in a nucleic acid population,

wherein the tagged nucleic acid probe has the structure

10

15

20

25

30

35

40

45

50

5

T^{ms}-L-X'

wherein.

T^{ms} is an organic group detectable by mass spectrometry, comprising carbon, at least one of hydrogen and fluoride, and optional atoms selected from oxygen, nitrogen, sulfur, phosphorus and iodine;

L is an organic group which allows a unique T^{ms}-containing moiety to be cleaved from the remainder of the compound, wherein the T^{ms}-containing moiety comprises a functional group which supports a single ionized charge state when the compound is subjected to mass spectrometry and is tertiary amine, quaternary amine or organic acid; and

X' is nucleic acid.

- 24. A method for determining the quantity of a single nucleotide polymorphism in a nucleic acid target comprising:
 - a) amplifying a sequence of a nucleic acid target containing a single nucleotide polymorphism;
 - b) generating a single strand form of the target;
 - c) combining a tagged nucleic acid probe with the amplified target nucleic acid molecules under conditions
 and for a time sufficient to permit hybridization of said tagged nucleic acid probe to complementary amplified
 selected target nucleic acid molecules, wherein said tag is correlative with a particular single nucleotide polymorphism and is detectable by spectrometry or potentiometry;
 - d) separating unhybridized tagged probe from hybridized tagged probe by a sizing methodology;
 - e) cleaving said tag from said probes;
 - f) detecting said tags by spectrometry or potentiometry; and
 - j) taking the ratio of tag correlated to the wild type polymorphism to the tag correlated with the mutant polymorphism, and determining therefrom the quantity of said polymorphism,

wherein the tagged nucleic acid has the structure

T^{ms}-L-X

wherein,

T^{ms} is an organic group detectable by mass spectrometry, comprising carbon, at least one of hydrogen and fluoride, and optional atoms selected from oxygen, nitrogen, sulfur, phosphorus and iodine;

L is an organic group which allows a unique T^{ms}-containing moiety to be cleaved from the remainder of the compound, wherein the T^{ms}-containing moiety comprises a functional group which supports a single ionized charge state when the compound is subjected to mass spectrometry and is tertiary amine, quaternary amine or organic acid; and

X' is nucleic acid.

Revendications

1. Composé de formule :

T^{ms}-L-X